

FP2027.ST25.txt
SEQUENCE LISTING

<110> Ruan, Yijun
Ng, Patrick
Wei, Chialin

<120> Gene identification signature (GIS) analysis

<130> FP2027

<160> 18

<170> PatentIn version 3.2

<210> 1

<211> 33

<212> DNA

<213> artificial

<220>

<223> GsuI-oligo dT primer

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<221> misc_feature

<222> (32)..(32)

<223> nucleotide v is

<220>

<221> misc_feature

<222> (32)..(32)

<223> nucleotide v is g, c or a

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<221> misc_feature

<222> (33)..(33)

<223> n is a, c, g, or t

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33

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<222> (5)..(38)

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<223> n is a, c, g, or t

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<223> primer Mme_mut1 used for the construction of vector pGIS1

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<223> primer Mme_mut2 used for the construction of vector pGIS1

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<222> (15)..(704)
<223> region from 15 to 704 represents the stuffer fragment, comprising
MmeI and BamHI, that is removed during cloning

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